



SEQUENCE LISTING

<110> Kellenberger, Johannes
Leadlay, Peter
Stauton, James
Stutzman-Engwall, Kim
McArthur, Hamish

<120> Polyketides, Their Perparation, and
Materials for Use Therein

<130> 0380-P02380US

<140> 09/743,162

<141> 2001-08-24

<150> PCT/GB99/02158

<151> 1999-07-06

<150> GB 9814622.8

<151> 1998-07-06

<160> 61

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 12381

<212> DNA

<213> Streptomyces avermitilis

<400> 1

cccgggcat	ctcccggatc	acctgtgcgg	ggctgggcat	gtgcaggaga	cactccaggg	60
cccacgccgc	gtcgaaggac	ccgtcgggaa	acggcagttc	catcgcgtcg	gcacgggtga	120
acacgacccg	gtccgccacg	tgcgactgct	tcgcgagagc	ggtcgccagc	ccgacctgaa	180
cctcgctcac	cgtcacgccg	acgacatcga	cgggcgcgct	cagggcgagc	cgcaccgccg	240
gctttccgga	accgcagccg	acgtccagga	cccggcggcc	cgtgatgcct	ctcagcttgc	300
cgatgaggag	atcggtgagc	cggtcggcgg	ccttgcccgg	tgaactgccg	tcccccggt	360
gcggccagta	tccgaggtgg	gtgttcccac	ccagcgcacg	attcatgagg	tcggtcaaac	420
ggtcgtagta	gtccccact	tccagggag	agggcgggg	ctgctccggg	acggccatca	480
tggtcgggaa	cctccgcaat	ccgggcgggg	cggcccagct	gtcgtggcga	tctactccag	540
gaaacgtcga	cctttttctg	ccacttgtcc	gagctatgca	gacaccccga	tcccctaaga	600
aatgaacacc	cttggggaacg	gcacagccca	gggggtggata	gggggtattcg	ccgcgcgcgc	660
gcggtcatta	gctttgaaga	gttgaagacg	ttcaagacat	tgatgcccg	ccgtcagcgg	720
atattctcgcg	ctcctttcat	tcttcgacgc	tgcatcgcag	ctctcatcat	gtccgcacgg	780
ccgccgagca	ttgcctagcg	gtgaggacac	agctcaggtg	cagaggatgg	acggcgggga	840
agaaccccg	cctgcggcag	gggaggtcct	cggagtggcc	gacgaggcgg	acggcggcgt	900
cgtcttcggt	tttcccgggc	agggcccgc	atggccgggc	atgggaagg	aacttctcga	960
cgcttccgac	gtcttccggg	agagcgctcg	cgcctgcgaa	gccgcgttcg	cgccctacgt	1020
cgactggtcg	gtggagcagg	tgttgccgga	ctcgcgggac	gctcccgggc	tggaccgggt	1080
ggacgtcgtc	cagccgacct	tgttcgcgt	catgatctcc	ctggccgccc	tctggcgctc	1140
gcaaggggtc	gagccgtgcg	cggtgctggg	acacagcctg	ggcgagatcg	cggcagccca	1200
cgtctcggga	ggcctgtccc	tggccgacgc	cgcacgcgtg	gtgacgcttt	ggagccaggc	1260
acagaccacc	cttgccggga	ccggcgcgct	cgtctccgtc	gccgccacgc	cggatgagct	1320
cctgccccga	atcgctccgt	ggaccgagga	caaccgcggc	cggctcgccg	tcgcagccgt	1380
caacggaccc	cggagcacag	tcgtttccgg	tgcccgcgag	gccgtcgcgg	accttgggtg	1440
cgacctcacc	gccgcgcagg	tgccgcagcg	catgatcccg	gtggacgttc	ccgcccactc	1500
ccccctgatg	tacgccatcg	aggaacgggt	cgtcagcggc	ctgctgccc	tcacccacg	1560
cccctcccgc	atccccttcc	actcctcggt	gaccggcggc	cgcctcgaca	ccgcgcagct	1620
agacgcggcg	tactgggtacc	gcaacatgtc	gagcacggtc	cggttcgagc	ccgccgccc	1680
gctgcttctg	cagcaggggc	ccaagacgtt	cgtcgagatg	agccgcacc	cgggtgctgac	1740
catgggcctc	caggagctcg	ccgcggacct	gggcgacacc	accggcaccg	ccgacaccgt	1800

gatcatggggc	acgctgcgcc	gcggccaggg	caccctggac	cacttcctga	cgtctctcgc	1860
ccaactacgg	gggcatgggtg	agacgtcggc	gaccaccgtc	ctctcggcac	gcctgaccgc	1920
gctgtccccc	acgcagcagc	agtcgtgtct	cctggacctg	gtgcgcgccc	acaccatggc	1980
ggtgtgaac	gacgacggaa	acgagcgcac	cgcgtcggat	gccggcccat	cggcgagttt	2040
cgccccacctc	ggcttcgact	ccgtcatggg	tgtcgaactg	cgcaaccgcc	tcagcaaggc	2100
caagggcctg	cggttgcccc	tgacgtctat	cttcgaccac	accacgcggg	ccgcggtcgc	2160
cgcgcgcctt	cggaccgcgg	cgctcggcca	cctcgacgag	gacaccgcgc	ccgtaccgga	2220
ctcaccacagc	ggccacggag	gcacggcgagc	ggcggacgac	ccgatcgcca	tcacggcat	2280
ggcatgccgt	ttccccggcg	gagtcgggtc	ccogaaggac	ctgtgggagc	tgccccgcctc	2340
gggaggagac	gccatcgggc	cgttccccac	cgaccgcgga	tggccccacg	aacagcgtca	2400
cgccccaggac	cccacgcagc	ccggcacgtt	ctatccgcag	ggaggcgggt	tccttcacga	2460
cgcggcgcac	ttcgacgcgc	gcttcttcgg	aatcagtcga	cgtgaggcac	tggcgatgga	2520
tccgcagcag	cggctgctgc	tggagacgtc	ctgggaggcg	ttcgagcggg	cgggaatcga	2580
tccgctgtcg	gtacgcgggt	cccgtacggg	cgtcttcgcg	ggcgccctct	ccttcgacta	2640
cggcccgcg	atggacaccg	cgtcgtcggg	gggcgcgcgc	gacgtggagg	gccacatcct	2700
caccgggtacc	acgggcagcg	tcctgtcggg	ccgtatcgcc	tacagcttcg	ggctggaagg	2760
gccggcgatc	accgtggaca	cgggggtgctc	ggcategctc	gtgacgctgc	atctggcgtg	2820
ccagtcgctg	cggtcgggtg	agtgcacgct	cgcgctggcc	ggcggcgctc	cggtcatgtc	2880
caccctcggc	atgttcacgc	agttctcccg	gcagcgcggg	ctgtcgggtg	acggcagggtg	2940
caaggcgctac	tcggctgcag	ccgacggcac	cggctggggc	gagggcgctc	ggatgctgtt	3000
ggtggagcgg	ttgtcggatg	cgggtcgggt	ggggcatcgg	gtgctggcgg	tggtagcgcg	3060
cagtgcgggtc	aaccaggacg	gtgcgtcgaa	tgggctgacg	gcgccgaacg	gtccggctca	3120
ggagcgggtg	atccggcagg	cgttggcgaa	cgcgggggtg	tccgtggcgg	atgtggaatgt	3180
ggtggagggg	cacgggacgg	gcacgacgct	gggtgatcgc	atcgaggcac	aggcgattgct	3240
cgccacgtac	gggcagcggg	cgggtgacag	cccgctgtgg	ctggggtctc	tgaagtccaa	3300
catcgggcac	accatggctg	ccgcgggtgt	gggtggggtc	atcaagatgg	tgatggcgtt	3360
gcgggagggg	gtgttgccgc	ggacgttgca	tgtggatgag	ccgtcgccgc	aggtggactg	3420
gtccgcgggg	gcgggtgcggc	tgtcgtcggg	ggcggtgccg	tggccggggg	acgcggcagg	3480
gcgggttcgg	cgggcgggag	tgtcgtcgtt	cgggatcggc	ggcacgaatg	cgcattgtgat	3540
tttgaggagg	gcgcggcgcg	cggggggctg	tgttgccggg	ggtgggggtg	tggagggtgc	3600
tccgggtcctt	gccatttcgg	tggctgagtc	ggtggccgct	ccagtggctg	tgtctgcgcc	3660
ggtggctgag	tcgggtgccg	tgcgggtgcc	ggtgccggtt	cctgtgccgg	tgtcggctag	3720
gtctgaggct	gggttgccgg	cgcaggcgga	ggcgttgctg	cagtacgtgg	cagtcgggcc	3780
ggacgttttcg	cttgccgatg	tgggtgcggg	tctggcctgt	gggcgggctg	tgtcggagca	3840
tcgtgcggtc	gtcctggccg	cggaccgtga	ggagctgggt	caagggttgg	gggcgctggc	3900
ggcgggtgag	cggatgcgc	gggtgaccac	gggtcatgcg	ccgggtgggtg	accggggcgg	3960
tgtcgtcttc	ctgtttcccg	gacagggtgg	gcagtgggcc	gggatgggtg	tgcgtctgct	4020
cgcctcctct	ccgggtgttcg	cccggcggat	gcaggcgtgc	gaggaggctc	tggcgccgtg	4080
ggtggactgg	tctgtggtgg	acatcctgcg	ccgggacgcg	gggatgcgg	tgtgggagcg	4140
ggccgatgtg	gtccagcctg	tgtgtttcag	cgtcatgggt	tctttggctg	ctctgtggcg	4200
ttcctacggt	atcgaaccgc	acgcggtcct	tggccattcc	cagggcgaga	tcgcgcgcgc	4260
gcatgtgtgt	ggggcgctga	gcctgaaggga	cgcggcgaa	actgttgccg	tgcgagccg	4320
ggcgctggcc	cgtgtgcggg	gcccggggcg	catggcctca	gtgccgctgc	ctgccagga	4380
ggtggagcag	ctcattgggtg	agcgggtggc	ggggcggttg	tgggtggcgg	cggtaaacgg	4440
cccccgctcc	accgccgtct	cgggggatgc	cgaggcgggtg	gacgaggtgc	tggcgtagctg	4500
tgccggcacc	gggggtgcggg	cccggcggat	cccgtcgcac	tatgcctcgc	actgccccca	4560
tgtgcagccc	ctgcgggagg	agttgctgga	gctgctgggg	gacatcagcc	cgcagccgtc	4620
cggcgtgccg	ttctttctcca	cgggtggagg	cacctggctg	gacaccacaa	ccctggacgc	4680
cgcctactgg	taccgcaacc	tgcaccagcc	ggtccgtttc	agcgatgccg	tccaggccct	4740
ggcggatgac	ggacaccgcg	tcttcgtcga	agtcagcccc	cacccccacc	tcgtccccgc	4800
catcgaagac	accaccgaag	acaccgcgga	agacgtcacc	gcgatcggca	gcctccgcgc	4860
cggcgacaac	gacaccgcgc	gcttctctca	cgccttcgcc	cacaccata	ccaccggcat	4920
cggcacaccc	accacctggc	accaccacac	caccaccacc	cacaccacc	cccaccccc	4980
cacgcacctc	gacctgcccc	cctaccctct	ccaacaccag	cactactggc	tcgagagctc	5040
acagccgggt	gccggatccg	gttcgggtgc	cggtgccggg	tcgggtgccg	gttcggggcg	5100
ggcagggact	gcgggcggga	cggcagaggt	ggagtgcgcg	ttctgggacg	cgggtggccc	5160
ccaggacctg	gaaacggtcg	cgaccacact	cgccgtgccc	ccctccgcgc	gcctggacac	5220
ggtggtgccc	gcactctcgc	cctggcaccg	ccaccaaac	gaccaagccc	gcataaacac	5280
ctggacctac	caggaaacct	ggaaaccctc	caccctcccc	accaccacc	aaaccaccca	5340
aacctggctc	atcgccatcc	ccgaaaccac	gaccaccac	ccccacatca	ccaacatcct	5400
caccaacctc	caccaccacg	gcataccccc	catccccctc	accctcaacc	acaccacac	5460
caacccccaa	cacctccacc	acaccctcca	ccacaccgga	caacaagccc	aaaaccacac	5520
caccggagcc	atcaccggcc	tgtctctcct	cctcgccctc	gacgaaacac	cccaccccc	5580
ccacccccac	acaccaccgc	gcaccctcct	caacctcacc	ctcaccacaa	cccacaccca	5640

aaccacccca	ccaaccccc	tctggtacgc	caccaccaac	gccaccacca	cccaccccaa	5700
cgacccccct	acacacccca	cccaagccca	aacctgggga	ctcgccccga	ccaccctcct	5760
cgaacacccc	accacacccg	ccggaatcat	cgacctcccc	accaccccca	ccccccacac	5820
cctccaccac	ctcacccaaa	ccctcaccca	accccaccac	caaacccaac	tcgccatccg	5880
caccacccgg	accacacccc	gccgcctcac	ccccaccacc	ctcaccceca	cacaccaacc	5940
acccaccccc	acccccacg	gaaccacct	catcacccgg	ggaaccggcg	ccctcgccac	6000
ccacctcacc	caccacctca	ccaccacca	acccacccaa	cacctcctcc	tcaccagccg	6060
aaccggcccc	cacaccccc	acgcacaaca	cctcaccacc	caactccaac	aaaaaggcat	6120
ccacctcacc	atcaccacct	gcgacaccag	caaccacagac	caactccaac	atctcctcaa	6180
caccatcccc	ccacaacacc	ccctcaccac	cgctcatccac	accgcaggca	tcctcgacga	6240
cggcaccctc	accaacctca	cccccaccca	actcaacaac	gtcctccgcg	ccaaagccca	6300
cagcgcccc	ctcctccacc	aactcaccca	acacaccccc	ctcacgcgct	tcgtcctcta	6360
ctcctccgcc	gccgccacct	tcggcgccac	cggccaagcc	aactacgccc	cagccaacgc	6420
ctacctcgac	gccctcgccc	accaccgcca	caccacccac	ctccccgcca	ccagcatcgc	6480
ctggggcacc	tggcaaggaa	acggactcgc	tgattcggac	aaggcccgcg	catatctcga	6540
ccgcgcgggg	tttcgaccga	tgtcacccga	gttggccacg	gcagcgggtca	cgcaggcgat	6600
cgcggacacc	gaacggccgt	atgtcgtcat	cgcgcacatc	gactggagca	agatcgaaca	6660
cacctctcag	accagaccgc	tggtgagcgc	ggcccgggaa	agggagccag	ctgtccagcg	6720
ccccactcca	ccggcgaggt	tgcacaaaac	gctggcccat	cagacgtcgg	ccgaccaacg	6780
ggccgcattg	ctcgagctcg	tacgagacca	tgtggcgcca	gtgctccggc	acgcggaccc	6840
gaaagccatc	gcgcccgcac	agtcgttccg	tgcactcggc	ttcgattcac	tcacggccgt	6900
cgagttccga	aacctgctga	tcaaggcaac	aggactccgc	cttcctgtct	cgctggtctt	6960
cgaccacccg	acccctgcca	aactcgccgt	acacctgcag	aaccaactgc	ggggcacagc	7020
agcgagtcg	gctccttcag	cggcagccgt	taccgcgag	gcttctgtca	ccgagccgat	7080
cgccatcggt	ggcatggcct	gtcgtttccc	cggcgagtg	acctcgccgg	acgacttctg	7140
ggatctgate	tcctccgagc	aggacgcgat	cggcgatttc	cccaccgacc	gcggctggga	7200
cctggacacg	ctctacgacc	ccgaccccca	cccccgcggc	acctgctaca	cccgaacccg	7260
cggattccct	tacgacgcag	gccacttcga	cgcgcgaattc	ttcggcatca	gcccccgcca	7320
agccctcgcc	atggaccccc	agcaacgact	cctcctcgaa	accgcctggg	aaaccatcga	7380
acacgcggcg	atcaaccccc	acaccctcca	cggcaccccc	accggagtct	tcaccggcac	7440
caacggacag	gactacgcac	ttcgcgtgca	caacgcgggc	cagtcaaccg	atggtttcgc	7500
actgaccgga	accgcgcgga	gcgtcatctc	cggctcgtatc	tcgtacacgt	ttggttttga	7560
gggtcctgcg	gtgtcggtgg	acacggcttg	ttcctcgctg	ttggtggctt	tgcattctggc	7620
ctgtcaggcg	ttgcgtgcgg	gtgagtgctc	gatggcgctt	gccggggggtg	tgacggtgat	7680
ttcgtctccg	ggtgccttcg	tggagttttc	gcggcagcgg	ggtctggccg	cggacgggca	7740
ttgcaaggcg	ttctcggcgg	cggcgagccg	cagccgctgg	ggtgaggggtg	tggggatgct	7800
gctggtggag	cggctctccg	acgcccacgc	caacgggtcac	cgtgtcctgg	ccgtggtgcg	7860
tggcagtgcg	gtcaaccagg	acggtgcgag	caacgggtctg	accgcgcccc	acgggcccgc	7920
ccagcagcgt	gtcatccgcc	aggccctcgc	caacgcgcggc	ttgtcggccg	gtgatgtcga	7980
cgcggtggag	gcccacggca	ccggcaccac	tttggggcgac	ccgatcgagg	cccaggccct	8040
cctcgcgacg	tacggacagg	accgtgcggg	cgaaggggcg	ctgtggctgg	gctcggctcaa	8100
gtccaatgtc	ggtcacacac	aggctgcgcg	gggcgtcgcc	ggggtgatca	agatggtgat	8160
ggcgtgcggg	catggtctgc	tgcgcgggac	gttgcatgtg	gatgagccgt	cgcgcgatgt	8220
ggactggtcc	gcgggtgcgg	tgcagctgct	gacggagacg	gtgccctggc	ccggcgggga	8280
ggggcgggcta	cggcgggag	gagtgctcatc	attcggcgctc	agcggcacca	acgcccacgt	8340
catcctcgaa	gaagcaccgc	ccgacgacgt	tcggggggga	ccaccgcggc	gcgaggggtga	8400
cgcgggcagc	gacgatgagg	ctgctgccgg	cagtccctggg	gtgtggccgt	ggctggtgtc	8460
ggccaagtgc	cagccggccc	tgcgcgcccc	ggcccaggcc	ctgcacgccc	acctcaccga	8520
ccaccccgcc	ctcgacctcg	cggatgtcgg	atacacccctc	gcccacgccc	gcgcctgttt	8580
cgaccaccgc	gccaccctca	tcgccgcgga	ccgcgacacg	ttcctgcaag	cactccaggc	8640
actcgccgca	ggcgagcccc	acccgcgcgt	catccacagc	agcgcgcccg	gcgggaccgg	8700
gaccggggag	gcccgcaggaa	agaccgcatt	cctctgctcc	ggacagggca	cccaacgccc	8760
cggcatggcc	cacggcctct	accacaccca	ccccgtcttc	gccgcgcgac	tcaacgacat	8820
ctgcacccac	ctcgaccccc	acctcgacca	ccccctcctc	ccccctcctca	cccaaaacga	8880
caacgacaac	gaggacgcgg	ccgactgtct	ccagcagacc	cgctacgccc	agcccgccct	8940
cttcgccttc	caggtcgccc	tccaccgcct	cctcaccgac	ggctaccaca	tcacccccca	9000
ctactacgcc	ggacactccc	tcggcgaaat	caccgcgcgc	cacctcgccg	gcactcctcac	9060
cctcaccgac	gccaccaccc	tcatacccca	acgcgccacc	ctcatgcaaa	ccatgccecc	9120
cggcaccatg	acaccacctc	acaccacccc	acccaccatc	acccaccacc	tcaccgccc	9180
cgaaaaacgac	ctcgccatcg	ccgccatcaa	cacccccacc	tcctctgtca	tcagcggcac	9240
ccccacaccc	gtccaacaca	tcaccacct	ctgccaacaa	caaggcatca	aaaccaaacc	9300
cctccccacc	aaccacgcct	tccactcccc	ccacaccaac	ccatcctca	accaactcca	9360
ccagcacacc	caaaccctca	cctaccaccc	acccacaccc	cccctcatca	ccgacaacac	9420
cccacccgac	caactcctca	ccccccacta	ctggacccaa	caagcccgca	acaccgtcga	9480

ctacgccacc	accacccaaa	ccctccacca	acacggcgtc	accacctaca	tcgatctcgg	9540
acccgacaac	accctcacca	ccctcaccca	ccacaacctc	cccaacaccc	ccaccaccac	9600
cctcacccctc	accaccccc	accaccaccc	ccaaacccac	ctcctcacca	acctcgccaa	9660
aaccaccacc	acctggcacc	cccaccacta	caccacccac	cacaaccaac	cccacacca	9720
caccacacctc	gacctcccca	cctaccctt	ccaacaccac	cactactggc	tcgaaagcac	9780
acagcccggt	gccggcaacg	tgtcagcagc	cggactcgac	cccaccgaac	acccctact	9840
cggcgccaca	ttggaactgg	cgactgacgg	tggagcgctt	cttgcagggc	gcttgtcttt	9900
gaggtcgcat	cogtggctgg	ctgaccatgc	cgtcggcggc	acggtgctgc	tgtcgggcgc	9960
caccttcctc	gaactcgccc	ttcatgcggg	cacatacgtg	ggctgcgacc	gagtggatga	10020
gctgacgctg	catgcgccc	tgggtggttc	tgtggatggg	ggtgtgagt	tgcaggttgg	10080
ggttgcggct	gcggatgggg	aggggcccgc	tttggtagt	gtgtatgcgc	ggggtgggag	10140
tgcttgtggt	gggggtggtg	cgtcgggtgg	ggtgtggacg	tgtcatgcct	cgggggtgct	10200
ggttgaggt	gctgctggtg	gtgtggtggt	ggatggtctg	gcgggggtgt	ggcgcgcgc	10260
gggtgcgggtg	gcggtggatg	tcgatggtgt	cgtgaccgt	ttggctgggg	ctggttgtgt	10320
tttggggccg	gtgttttcgg	ggctgcgtgc	ggtgtggcgt	gatggggggg	atttgctggc	10380
tgaggtgtgt	ctgcggagg	aggcgtgggg	tgatgcggct	ggttttgggc	tgcacccggc	10440
ggtgctggat	ggtgtggtcc	agccgttgct	ggtgttgctt	ccgggtggga	cggggtttgg	10500
ggagggggcg	gggttcgggg	aggtgttccg	ggtgccggct	gtgtggggtg	gtgtgtcgt	10560
tcaccggggc	ggtgtgaccg	gtgtgcgggt	gcgtgtgtcg	gctgtcgggc	ggggcggcgc	10620
gcgtgaggcg	gtgtcggctc	tggtcgggga	tgaggcgggt	gtgccggtgg	cgtcggtcga	10680
tcgtcttgag	ttgcggcctg	tggatatggg	tcagttgcgt	gctgtctcgg	tttcggcggg	10740
gcggcgggggt	tcgctgtatg	cgggtgcagt	ggctgaggtg	ggtcctgtgc	cgggtgtgtg	10800
gcaggcgtgg	gcgtggcacg	aggacgtggg	tgagagcggg	ggtgggcctg	tgccgggggt	10860
ggtgggtggtg	cgggtcccgg	atgccgggtc	cggtgccggg	ggcgggtggc	gtggtggcgg	10920
tggtgtgggt	gaggttgttg	gtggggtgtt	gggtgtggtg	caggggtggc	tggggctgga	10980
gcggtttgcg	ggttcgcggc	tgggtggtgt	gacccggggg	gcggtggtgg	ccggcccggg	11040
ggacggcccc	gtggatgtgg	tgggtgcgtc	ggtgtggggg	ctggtgcgtt	cggcgcaggc	11100
tgagcatccg	gaccggtttg	tcctcctcga	cctcgacacc	gacaccggca	ccgacctcga	11160
caccggtgct	ggtgctggtt	ggggcgtgga	tgggtggcgt	gtggcggcgg	tgggtggcgtg	11220
tgggtgcggc	cagttggcgg	tgcgtgggga	gcggttgctg	gccgcacgcc	tgacacgact	11280
tgagtcaccc	ggtgatgttc	cagcccagcg	gtccggtgac	acacgagccc	ggcgggtccga	11340
cgtgcctgcc	cagcgtcccg	gtggcgtgcc	tgtcggcgg	tcggttgatg	tatcgggtcg	11400
ggaggtggtg	ccgtggttgt	cgggtgggtc	ggtgttggtg	acgggtggga	cgggtgtgct	11460
gggtgcggcg	gtggcgcggc	atctggctgg	tgtgtgtggg	gtgcgggatc	tgtgttgtgt	11520
gagccggcgt	ggtccggatg	ctccgggtgc	ggaggggtctg	cgggcggagc	tggcccggtt	11580
ggggcgagg	gtgcggattg	ttgcgtgtga	tgtgggggag	cggcgggagg	tgggtccggt	11640
gctggagggt	gttcctgccg	ggtgtccgct	gacgggtgtc	gtgcatgcgg	ctggtgtgct	11700
ggacgatgcg	acgatgcct	ctctcacgcc	cgagcggctg	ggcacggtgt	tcgcggccaa	11760
ggtggatgcc	gctcttttgc	tggatgagct	gacgcggggg	atggagctgt	cggcgttcgt	11820
gctgttctcc	tcggccgcgg	ggatcctggg	gtcggccggg	cagggcaact	acgccgcggc	11880
caatgccgct	ctggacgcgc	tggcgtaccg	gcggcggg	gcgggtctgc	cgggggtgtc	11940
gctggcggtg	gggctgtggg	aagaggccag	cgggatgacc	gggcacctgg	ccggcaccca	12000
ccaccggcgc	atcatccgtt	cgggtctgca	tcccatgtcg	accccgagcg	cactggccct	12060
cttcgatgcg	gccctggctc	tggaccggcc	ggtcctgctg	cccgcggacc	tgcgtccgc	12120
cccgcctctg	ccgcccctgc	tgcaggacct	cctgcccgc	acccgcggc	gcaccacccg	12180
caccaccact	accggtggtg	cggacaacgg	cgcccagctg	cacgcccggc	tggccggcca	12240
gacacagcaa	caacagcaca	ccacctcct	cgccctggct	cgctccaca	tcgccaccgt	12300
cctggggccac	accacccccg	acaccatccc	ccccgaccgc	gcgttccgcg	acctcggtt	12360
cgactccctc	accgccgtcg	a				12381

<210> 2
 <211> 37
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 2
 tacctaggcc gggccggact ggtcgacctg ccgggtt

37

<210> 3
 <211> 30
 <212> DNA

<213> Artificial Sequence
 <220>
 <223> Primer
 <400> 3
 atgttaaccg gtcgcgagg ctctccgtct 30
 <210> 4
 <211> 32
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 4
 atgttaaccg gtctgcccgc tgccgagcgg ac 32
 <210> 5
 <211> 30
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 5
 cttctagact atgaattccc tccgcccagg 30
 <210> 6
 <211> 28
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 6
 atactagtcc tcgtgacgag ctgcacgg 28
 <210> 7
 <211> 30
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 7
 taatgcatcc ggttctccgg cccgctcgct 30
 <210> 8
 <211> 28
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 8
 taagatcttc cgacctacgc cttccaac 28
 <210> 9

<211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 9
 taatgcatcg acctcgttgc gtgccgcggt 30

 <210> 10
 <211> 39
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 10
 atagatctgc ctacgtaccc gttcgaacac cagcgcttc 39

 <210> 11
 <211> 50
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 11
 atcctcaggt tcggccctgc cgctcggcc tgcgcggcgg cgcgcagctt 50

 <210> 12
 <211> 28
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 12
 taagatcttc cgacgtacgc gttccagc 28

 <210> 13
 <211> 33
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 13
 atgctagcca ctgcgccgac gaatcaccgg tgg 33

 <210> 14
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 14
 taagatctcc ctacgtaccc cttcaaccac 30

<210> 15
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 15
 gctagccgcc gcgccagctc gggc 24

 <210> 16
 <211> 29
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 16
 cctacgtacg ccttcgacca ccagcactt 29

 <210> 17
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 17
 cggctagcgg gcgttccagg ccgccgtcct 30

 <210> 18
 <211> 34
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 18
 tacctgaggg accggctagc gggctctgccg cgtg 34

 <210> 19
 <211> 34
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 19
 atgctagccg ttgtgccggc tcgccggtcg gtcc 34

 <210> 20
 <211> 34
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

<400> 20	
gagcccggaat tcttcggcat cagccccgc gaag	34
<210> 21	
<211> 63	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 21	
gagctagcag gtggggagat ctaggtgggt gtgggtgtgg ggttggttgt ggtgggtgggt	60
gta	63
<210> 22	
<211> 34	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 22	
gccccgctag ccggccagac acacgaacaa cagc	34
<210> 23	
<211> 34	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 23	
gggaattcct cgaggatgac gtgggcgttg gtgc	34
<210> 24	
<211> 30	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 24	
taagatctag cgctccgagg ttcttgcccg	30
<210> 25	
<211> 30	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 25	
atgctagcct accgctgccc gggccgccc	30
<210> 26	
<211> 34	
<212> DNA	
<213> Artificial Sequence	

<220>
 <223> Primer

 <400> 26
 cctagatccg cccacctacc ccttccaaca ccag 34

 <210> 27
 <211> 36
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 27
 tgggctagcg ttttgtgcaa ctccgccggt ggagtg 36

 <210> 28
 <211> 37
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 28
 tggctgcaga gctcacagcc gggtgccgga tccggtt 37

 <210> 29
 <211> 36
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 29
 tttcctcagg tccgccggtg gagggggag ctggac 36

 <210> 30
 <211> 42
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 30
 cctagatctc cccacctacc ccttccaaca ccaccactac tg 42

 <210> 31
 <211> 40
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Primer

 <400> 31
 ccggctagcc gggcgtagc ctggggcgcc ttgtccgcac 40

 <210> 32
 <211> 39
 <212> DNA

<213> Artificial Sequence
 <220>
 <223> Primer
 <400> 32
 ccctacgtac cccttccaac accactactg gctcgaaag 39
 <210> 33
 <211> 37
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 33
 ggccctcagg tggcgccgt tgtccgcacc accggta 37
 <210> 34
 <211> 27
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 34
 tacctaggca ccaccacaac ccgggta 27
 <210> 35
 <211> 28
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Primer
 <400> 35
 tacaattggc ccgcgagtcc ccgacgct 28
 <210> 36
 <211> 35
 <212> PRT
 <213> Streptomyces erythraea
 <400> 36
 Val Ala Val Asp Trp Glu Ala Val Leu Gly Arg Ala Gly Leu Val Asp
 1 5 10 15
 Leu Pro Gly Tyr Pro Phe Gln Gly Lys Arg Phe Trp Leu Leu Pro Asp
 20 25 30
 Arg Thr Thr
 35
 <210> 37
 <211> 35
 <212> PRT
 <213> Streptomyces erythraea
 <400> 37
 Val Thr Thr Ala Pro Ser Glu Arg Ala Gly Glu Pro Glu Thr Glu Ser
 1 5 10 15
 Leu Arg Asp Arg Leu Ala Gly Leu Pro Arg Ala Glu Arg Thr Ala Glu
 20 25 30

Leu Val Arg
 35

 <210> 38
 <211> 41
 <212> PRT
 <213> Streptomyces erythraea

 <400> 38
 Val Ala Val Asp Trp Glu Ser Val His Leu Gly Thr Gly Ala Arg Arg
 1 5 10 15
 Val Pro Leu Pro Thr Tyr Pro Phe Gln Arg Glu Arg Val Trp Leu Glu
 20 25 30
 Pro Lys Pro Val Ala Arg Arg Ser Thr
 35 40

 <210> 39
 <211> 33
 <212> PRT
 <213> Streptomyces erythraea

 <400> 39
 Asp Asp Ala Arg Arg Ala Ala Pro Gly Ala Pro Ala Glu Pro Arg Val
 1 5 10 15
 Gly Ala Leu Ala Ser Leu Pro Ala Pro Glu Arg Glu Glu Ala Leu Phe
 20 25 30
 Glu

 <210> 40
 <211> 41
 <212> PRT
 <213> Streptomyces erythraea

 <400> 40
 Val Gly Ala Asp Leu Arg Pro Ala Val Ala Gly Gly Arg Pro Ala Glu
 1 5 10 15
 Leu Pro Thr Tyr Pro Phe Glu His Gly Arg Phe Trp Pro Arg Pro His
 20 25 30
 Arg Pro Ala Asp Val Ser Ala Leu Gly
 35 40
 \
 <210> 41
 <211> 35
 <212> PRT
 <213> Streptomyces erythraea

 <400> 41
 Arg Ala Lys Leu Arg Ala Ala Gly Gly Ala Glu Ala Ala Gly Pro Asn
 1 5 10 15
 Val Val Asp Arg Leu Ala Gly Arg Ser Glu Ser Asp Gln Val Ala Gly
 20 25 30
 Leu Ala Glu
 35

 <210> 42
 <211> 38
 <212> PRT
 <213> Streptomyces erythraea

 <400> 42
 Val Glu Val Asp Trp Ser Pro Ala Phe Ala Asp Ala Arg Pro Val Glu
 1 5 10 15
 Leu Pro Val Tyr Pro Phe Gln Arg Gln Arg Tyr Trp Leu Pro Ile Pro

Thr Gly Gly Arg Ala Arg 20 25 30
 35
 <210> 43
 <211> 34
 <212> PRT
 <213> Streptomyces erythraea
 <400> 43
 Ala Gly Ala Arg Ala Glu Ala Arg Gln Ser Glu Glu Gly Pro Ala Leu
 1 5 10 15
 Ala Gln Arg Leu Ala Ala Leu Ser Thr Ala Glu Arg Arg Glu His Leu
 20 25 30
 Ala His
 <210> 44
 <211> 40
 <212> PRT
 <213> Streptomyces hygroscopicus
 <400> 44
 Val Thr Val Asp Trp Pro Ala Ile Leu Gly Thr Thr Thr Ala Arg Val
 1 5 10 15
 Leu Asp Leu Pro Thr Tyr Ala Phe Gln His Gln Arg Tyr Trp Val Lys
 20 25 30
 Ser Val Asp Arg Ala Ala Ala Asp
 35 40
 <210> 45
 <211> 32
 <212> PRT
 <213> Streptomyces hygroscopicus
 <400> 45
 Arg Pro Ile Ala Arg Arg Ala Ala Ser Thr Gly Asp Ser Ser Val Gln
 1 5 10 15
 Trp Leu Ala Ala Leu Ala Pro Glu Glu Arg Ala Lys Ala Leu Leu Arg
 20 25 30
 <210> 46
 <211> 40
 <212> PRT
 <213> Streptomyces hygroscopicus
 <400> 46
 Val Thr Val Asp Trp Pro Ala Ile Leu Gly Thr Ala Thr Thr Arg Val
 1 5 10 15
 Pro Asp Leu Pro Thr Tyr Ala Phe Gln His Gln Arg Phe Trp Ala Glu
 20 25 30
 Gly Ala Asp Arg Ser Val Ala Gly
 35 40
 <210> 47
 <211> 32
 <212> PRT
 <213> Streptomyces hygroscopicus
 <400> 47
 Arg Pro Val Ala Arg Arg Ala Ala Ser Thr Gly Gly Ser Ser Val Gln
 1 5 10 15
 Trp Leu Ala Arg Leu Ala Pro Val Glu Arg Glu Lys Ala Leu Leu Lys
 20 25 30

<210> 48
 <211> 44
 <212> PRT
 <213> Streptomyces hygrosopicus

 <400> 48
 Val Thr Val Asp Trp Arg Ala Val Leu Gly Asp Val Pro Ala Thr Arg
 1 5 10 15
 Val Leu Asp Leu Pro Thr Tyr Ala Phe Gln His Gln Arg Tyr Trp Ala
 20 25 30
 Glu Ala Gly Arg Ser Ala Asp Val Ser Ala Ala Gly
 35 40

 <210> 49
 <211> 32
 <212> PRT
 <213> Streptomyces hygrosopicus

 <400> 49
 Arg Pro Val Ala Arg Arg Ala Ala Ser Thr Gly Asp Ser Ser Ala Gln
 1 5 10 15
 Trp Leu Val Gly Leu Ala Pro Glu Glu Arg Ala Lys Ala Leu Leu Lys
 20 25 30

 <210> 50
 <211> 40
 <212> PRT
 <213> Streptomyces hygrosopicus

 <400> 50
 Val Thr Val Asp Trp Pro Ala Ile Leu Gly Thr Thr Thr Thr Arg Val
 1 5 10 15
 Leu Asp Leu Pro Thr Tyr Ala Phe Gln His Gln Arg Tyr Trp Leu Lys
 20 25 30
 Ser Val Asp Arg Ala Ala Ala Asp
 35 40

 <210> 51
 <211> 35
 <212> PRT
 <213> Streptomyces hygrosopicus

 <400> 51
 Arg Pro Gln Ser Arg Thr Ala Ala Arg Asn Glu Val Gly Ser Gln Pro
 1 5 10 15
 Leu Ser Ala Arg Leu Thr Gly Arg Thr Ser Val Glu Gln His Arg Ile
 20 25 30
 Met Leu Glu
 35

 <210> 52
 <211> 24
 <212> PRT
 <213> Streptomyces avermitilis

 <400> 52
 Thr His Pro His Pro His Thr His Leu Asp Leu Pro Thr Tyr Pro Phe
 1 5 10 15
 Gln His Gln His Tyr Trp Leu Glu
 20

 <210> 53
 <211> 25

```

<212> PRT
<213> Streptomyces avermitilis

<400> 53
Pro Thr Pro Pro Ala Glu Leu His Lys Thr Leu Ala His Gln Thr Ser
1      5      10      15
Ala Asp Gln Arg Ala Ala Leu Leu Glu
20      25

<210> 54
<211> 24
<212> PRT
<213> Streptomyces avermitilis

<400> 54
Asn Gly Pro His Thr His Thr His Leu Asp Leu Pro Thr Tyr Pro Phe
1      5      10      15
Gly His His His Tyr Trp Leu Glu
20

<210> 55
<211> 25
<212> PRT
<213> Streptomyces avermitilis

<400> 55
Ala Asp Asn Gly Ala Gly Leu His Ala Arg Leu Ala Gly Gln Thr His
1      5      10      15
Glu Gln Gly His Thr Thr Leu Leu Ala
20      25

<210> 56
<211> 122
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 56
ctaggccggg ccggactggg agatctgcct acgtatcctt tccagggcaa gcggttctgg      60
ctgcagccgg accgcactag tcctcgtgac gagggagatg catcgagcct gagggaccgg      120
tt                                                    122

<210> 57
<211> 118
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

<400> 57
aaccggtccc tcaggctcga tgcattctccc tcgtcacgag gactagtgcg gtccggctgc      60
agccagaacc gcttgccctg gaaaggatac gtaggcagat ctaccagtcc ggcccggc      118

<210> 58
<211> 66
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide

```

<400> 58
 ctaggccggg ccggactggt agatctgcct acgtatcctt tccagggcaa gcggttctgg 60
 ctgcag 66

<210> 59
 <211> 62
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide

<400> 59
 ctgcagccag aaccgcttgc cctggaaagg atacgtaggc agatctacca gtccggccccg 60
 gc 62

<210> 60
 <211> 56
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide

<400> 60
 ccggaccgca ctagtctctg tgacgagggg gatgcatcga gcctgagggg ccggtt 56

<210> 61
 <211> 56
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide

<400> 61
 aaccggtccc tcaggctcga tgcattctccc tcgtcacgag gactagtgcg gtccgg 56